

rerun

#14/ K.T. 1600
12/31/01RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/486,757DATE: 12/27/2001
TIME: 11:30:01Input Set : A:\J&J1673seqlist.txt
Output Set: N:\CRF3\12272001\I486757.rawRaw
Seq
Listing

3 <110> APPLICANT: Kutchan, Toni
4 Zenk, Meinhart
5 Atkins, David
7 <120> TITLE OF INVENTION: Cytochrome P450 Reductases from Poppy Plants
9 <130> FILE REFERENCE: J&J 1673
11 <140> CURRENT APPLICATION NUMBER: US 09/486757
C--> 12 <141> CURRENT FILING DATE: 2000-02-28
14 <150> PRIOR APPLICATION NUMBER: AU.P08872
15 <151> PRIOR FILING DATE: 1997-08-29
17 <160> NUMBER OF SEQ ID NOS: 32
19 <170> SOFTWARE: PatentIn version 3.0
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40 Glu Lys
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56 <211> LENGTH: 8
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ENTERED

see page 5

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Input Set : A:\J&J1673seqlist.txt

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124 tcagaatata tttctgaccc aattttcatt atggtcacaa ctgtagcttc aatgctgatt      240
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130 ctactatat tttttggtac tcagactggg actgctgaag gatttgctaa ggcattggca      420
132 gaagaaatta aggcaaagta caagaaagca gttgttaaag tagttgacct ggatgactat      480
134 gcagccgagg atgatcaata tgaagagaaa ttaaagaaag agtctttggt gtttttcatg      540
136 gtagccactt atggtgatgg tgagccaact gacaatgctg cgagatttta caaatgggtc      600
138 actcaggaac atgaaagggg agagtggctt cagcaactaa cttatggtgt ttttggtttg      660
140 ggtaaccgtc aatacagaca tttcaacaag atcgcggtag atgtggatga gcaactcgtt      720
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158 aaagaagacg ggtcacccca gggaagctca ttaccacctc ctttcccagg tccttgcacc 1260
160 ttacgatctg ccttagcacg ctatgctgat cttttgaatc ctctagaaa ggcttctctg 1320
162 attgctctgt ccgctcatgc atctgtaccc agtgaagcag agagattgcg ctttttgtca 1380
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172 ggagtgtgtt cgacatggat gaagcatgca gttcctcagg atagctgggc tcctattttt 1680
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211 <212> TYPE: DNA

212 <213> ORGANISM: Eschscholzia californica

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220 aaaactatgg aacaaactgc ggttaaagtc tctttgtttg atctattttc ttcgatactt 180
222 aatggaaagt tggatccgtc gaacttttct tcagattcaa gtgctgctat tttgattgaa 240
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250 cgttcaacac ccatgcagag ccaatgtcgt tgtaaagaag gagcttcaca ctccagtatc 1080
252 tgatcgtttt tgtattcatc tgggaatttga catttctggc actgggctca cgtatgaaac 1140
254 aggagaccat gtcggtgttt actctgagaa ttgtgttgaa gttgtcgagg aagcagagag 1200
256 gctattgggt tactcatcag acaccgtttt ttcaatccat gtcgataaag aggacggctc 1260
258 acccattagt ggaagcgtc tagctcctcc ttttccaact cctgcactc taagaacagc 1320
260 actaacacga tacgtgatc tgttgaattc tccaagaag gctgctctgc atgctttggc 1380
262 tgcttatgca tccgatccaa aggaagcggg gcgactaagg tatcttgctg ctctgctgg 1440
264 gaaggacgaa tacgcccagt ggatagtagc tagtcagaga agtctgctag tggatcatgg 1500
266 tgaattccca tcagcaaagg ctccaattgg ggttttcttt gcagcagtag ctctcgtctt 1560
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316 tcagaatata tttctgaccc aattttcatt atggtcacaa ctgtagcttc aatgctgatt 240
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320 tataaaacaa taattgataa agaagaagag gagattgaag ttgactctgg taaaattaag 360
322 ctactatat tttttgttac tcagactggg actgctgaag gatttgctaa ggcattggca 420
324 gaagaaatta aggcaaagta caagaaagca gttgttaaag tagttgacct ggatgactat 480
326 gcagccgagg atgatcaata tgaagagaaa ttaaagaaag agtctttggg gtttttcatg 540
328 gtagccactt atgggtgatg tgagccaact gacaatgctg cgagatttta caaatggttc 600
330 actcaggaac atgaaagggg agagtggctt cagcaactaa cttatgggtg ttttggtttg 660
332 ggtaaccgtc aatacgagca tttcaacaag atcgcggtag atgtggatga gcaactcggg 720
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356 tcacctctgg gaaagaatga gtattcaaaa tgggtagttg gaagtcagag gagtcttttg 1440
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394 gatcttctct ttggattgat tactgtaagt tctaaccaga tgatagattg tacttaaaga 2580
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404 <213> ORGANISM: Papaver somniferum

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412 20 25 30
414 Thr Val Ala Ser Met Leu Ile Gly Phe Gly Phe Phe Ala Cys Met Lys
415 35 40 45
417 Ser Ser Ser Ser Gln Ser Lys Pro Ile Glu Thr Tyr Lys Pro Ile Ile
418 50 55 60
420 Asp Lys Glu Glu Glu Glu Ile Glu Val Asp Pro Gly Lys Ile Lys Leu
421 65 70 75 80
423 Thr Ile Phe Phe Gly Thr Gln Thr Gly Thr Ala Glu Gly Phe Ala Lys
424 85 90 95

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\J&J1673seqlist.txt

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L:35 M:283 W: Missing Blank Line separator, <400> field identifier
L:47 M:283 W: Missing Blank Line separator, <400> field identifier
L:59 M:283 W: Missing Blank Line separator, <400> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:95 M:283 W: Missing Blank Line separator, <400> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:116 M:283 W: Missing Blank Line separator, <400> field identifier
L:213 M:283 W: Missing Blank Line separator, <400> field identifier
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L:405 M:283 W: Missing Blank Line separator, <400> field identifier
L:542 M:283 W: Missing Blank Line separator, <400> field identifier
L:636 M:283 W: Missing Blank Line separator, <400> field identifier
L:775 M:283 W: Missing Blank Line separator, <400> field identifier
L:853 M:283 W: Missing Blank Line separator, <400> field identifier
L:989 M:283 W: Missing Blank Line separator, <400> field identifier
L:1069 M:283 W: Missing Blank Line separator, <400> field identifier
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L:1875 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1893 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1971 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:2031 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32